

GenCore version 6.3
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OM nucleic - nucleic search, using sw model

Run on: June 26, 2009, 03:19:14 ; Search time 612 Seconds
(without alignments)
14891.617 Million cell
updates/sec

Title: US-09-300-482-225
Perfect score: 316
Sequence: 1
gataagggtgcgcacactgag.....gaatggagggagtacttcga 316

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 41078765 seqs, 14427166270 residues

Total number of hits satisfying chosen parameters: 82157530

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published_Applications_NA_Main:*

1:
/ABSS/Data/CRF/ptodata/1/pubpna/US07_PUBCOMB.seq:*

2:
/ABSS/Data/CRF/ptodata/1/pubpna/US08_PUBCOMB.seq:*

3:
/ABSS/Data/CRF/ptodata/1/pubpna/US09A_PUBCOMB.seq:*

4:
/ABSS/Data/CRF/ptodata/1/pubpna/US09B_PUBCOMB.seq:*

5:
/ABSS/Data/CRF/ptodata/1/pubpna/US09C_PUBCOMB.seq:*

6:
/ABSS/Data/CRF/ptodata/1/pubpna/US09D_PUBCOMB.seq:*

7:
/ABSS/Data/CRF/ptodata/1/pubpna/US10A_PUBCOMB.seq:*